

1	GCACGAGGAACAGAACACTTCTCATGTCAGGGTCAGATTACAAGAGCACTCAAGACTT	60
61	TACTGAGCAGAAACTCAGGAATCCTCTATCAAAAGAGGTTGGCAACTAAACTAAGACA	120
121	TTAAAAGGAAATACCAGATGCCACTCTGCAGGCTGCAATAACTACTACTTACTGGATAC	180
181	ATTCAAACCCCTCCAGAACAGTTACAGGTAAACCAACAAGAAATGCAAGCGTCGAC	240
1	M Q A V D	5
241	AATCTCACCTCTGCGCCTGGGAACACCAGTCTGTGCAACCAGAGACTACAAAATCACCAG	300
6	N L T S A P G N T S L C T R D Y K I T Q	25
301	GTCCCTTCCCCACTGCTCTACACTGTCTGTGTTTTGTTGGACTTATCACAATGGCTG	360
26	V L F P L L Y T V L F F V G L I T N G L	45
361	GCGATGAGGATTTCTTCAAATCCGGAGTAAATCAAACTTATTATTTTCTTAAGAAC	420
46	A M R I F F Q I R S K S N F I I F L K N	65
421	ACAGTCATTTCTGATCTTCTCATGATTCTGACTTTCCATTCAAATCTTAGTGTGATGCC	480
66	T V I S D L L M I L T F P F K I L S D A	85
481	AAACTGGGAACAGGACCACTGAGAACCTTGTGTGTCAGTTACCTCCGTCAATT	540
86	K L G T G P L R T F V C Q V T S V I F Y	105
541	TTCACAATGTATATCAGTATTTCATTCCGGACTGATAACTATCGATCGCTACCAAGAG	600
106	F T M Y I S I S F L G L I T I D R Y Q K	125
601	AcCACCAGGCCATTAAAACATCCAACCCAAAAATCTTGGGGCTAAGATTCTCTCT	660
126	T T R P F K T S N P K N L L G A K I L S	145
661	GTTGTCATCTGGCATTCTCATGTTCTTACTCTCTTGCCTAACATGATTCTGACCAACAGG	720
146	V V I W A F M F L L S L P N M I L T N R	165
721	CAGCCGAGAGACAAGAATGTGAGAAATGCTCTTCTTAATCAGAGTTGGTCTAGTC	780
166	Q P R D K N V K K C S F L K S E F G L V	185
781	TGGCATGAAATAGTAAATTACATCTGCAAGTCATTCTGGATTAAATTCTTAATTGTT	840
186	W H E I V N Y I C Q V I F W I N F L I V	205

FIG. 1A

841	ATTGTATGTTATACACTCATTACAAAAGAACGTGACCGGTACACGTAAGAACGAGGGGT	900
206	I V C Y T L I T K E L Y R S Y V R T R G	225
901	GTAGGTAAAGTCCCCAGGAAAAAGGTGAACGTCAAAGTTTCATTATCATTGCTGTATTG	960
226	V G K V P R K K V N V K V F I I I A V F	245
961	TTTATTTGTTTGTCTTCCATTGCGGAATTCTTACACCTGAGCCAAACCCGG	1020
246	F I C F V P F H F A R I P Y T L S Q T R	265
1021	GATGTCTTGACTGCACTGcTGAAAATACTCTGTTCTATGTGAAAGAGAGGACTCTGTGG	1080
266	D V F D C T A E N T L L F Y V K E S T L W	285
1081	TTAACCTCTTAAATGCGATGCCCTGGATCCGTTCATCTATTTTCTTGCAGTCCTTC	1140
286	L T S L N A C L D P F I Y F F L C K S F	305
1141	AGAAATTCTTGTAAAGTGTGAAAGTGGCCCAATTCTGCAACATCTCTGTCCAGGAC	1200
306	R N S L I S M L K C P N S A T S L S Q D	325
1201	AATAGGAAAAAGAACAGGATGGGGTGAcCCAAATGAAGAGACTCCAATGTAACAAAT	1260
326	N R K K E Q D G G D P N E E T P M *	343
1261	TAACTAAGGAAATATTCAATCTCTTGTGTTAGACTCGTTAAAGCAAAGCGCTAAGT	1320
1321	AAAAATATTAACTGACGAAAGCAACTAAGTTAATAATGACTCTAAAGAAACAGAA	1380
1381	GATTACAAAAGCAATTTCATTACCTTCCAGTATGAAAAGCTATCTTAAATATAGAA	1440
1441	AACTAATCTAAACTGTAGCTGTATTAGCAGCAAAACAAACGACATCCAATTGTCATGCTG	1500
1501	CATGCAAAACTACACAGAATTCTGTTTGgCAGAGTTGGCAAAATGAGTAATCATAT	1560
1561	AATATTACTGTAATTAAAATACATTATGTTACAATTATTTTCTATAATCAA	1620
1621	CTAAGGAAGAACGATCAATTGGATATAATCTCTTACCAAAATGATAGTTAAATGTAT	1680
1681	ATATATCCTAGTCCCCTAACCAATCCTGACCTATTGGGATACTTATAAAAATTAGTA	1740
1741	AGTGGGATACACAAAGAACATAACTTAACTTTTCAATTAGCcAAAAACCTAAGGG	1800

FIG. 1B

Appl. No. To be assigned; Group Art Unit: To be assigned
Att. No. 1488.1220003/EKS/EJH.
Inventors: Li *et al.*; Tel: 202/371-2600
Title: Human G-Protein Coupled Receptors

1801 ATTTAACTAATTGAAaCTGTATTGATTGGACTTAATTTTTATGTTTATTAGAAGAT 1860
1861 AAAGATTTAAGAAGACCTTACAATAAAGAGAAGAAATATCGAAGTCATTAAAATAAGGA 1920
1921 GACTTACTTTATGACATTCTAATACTAAAAAAATAGAAATATTCCTTAATTCTAGAG 1980
1981 AAACTAGTTTACTAATTTTACAACCTCAATAATACCATCACTGACACTTACCTTAT 2040
2041 TAATTAGCTCTAGAAAATAGCTGCTAATTAGGTTAATGAACATTTACCTTAGTGAAAA 2100
2101 AAAaTTAATTAAATATGATTACAAAGTTGACAGCATAACTACTGAGAGGAAAGTGATTG 2160
2161 ATCTGTTGTAATTACTTGTGTTGTATTGGTGTATAAAATACAAATTACATTAACCTC 2220
2221 TAAAtcattaaaAAAAAAAAAAAAAA 2247

FIG. 1C

1 MQAVDNLTSAPI...GNTSLCTR DYKITQVLFPLLYTVLFFVGLITNGLA 46
 :|.:|.:|...||.:|...|...|.:|.:|.:| ||
 3 IQMANNFTPPSATPQGNDCDLYAHHSTARIVMPLHYSLVFIIGLVGNLLA 52
 :|.:|.:|...||.:|...|...|.:|.:|.:| ||
 47 MRIFFQIRSKNS.FIIFLKNTVISDLMILTFPFKILSDAKLGTGPLRTF 95
 :|.:|.:|...||.:|...|...|.:|.:|.:| ||. .: .
 53 LVVIVQNRKKINSTTLYSTNLVISDILFTALPTRIAYYAMGFDWRIGDA 102
 :|.:|.:|...||.:|...|...|.:|.:|.:| ||. .: .
 96 VCQVTSVIFYFTMYISISFLGLITIDRYQKTTTRPFKTSNPKNLLGAKILS 145
 :|.:|.:|...||.:|...|...|.:|.:|...|.:|.:| ||. .: .
 103 LCRITALVFYINTYAGVNFMTCCLSIDRFIAVHPLRYNKIKRIEHAKGVC 152
 :|.:|.:|...||.:|...|...|.:|.:|...|.:|.:| ||. .: .
 146 VVIWAFMFLLSLPNMI..LTNRQPRDKNVKKCSFLKSEFGLVWHEIVNYI 193
 :|.:|.:|...||.:|...|...|.:|...|...|.:|...|.:|...| ||. .: .
 153 IFWVILVFAQTLPLLIINPMISKQEAERITCMEYPNFEETKSLPWILLGACF 202
 :|.:|.:|...||.:|...|...|.:|...|...|.:|...|.:|...| ||. .: .
 194 CQVIFWINFLIVIVCYTLITKELYRSYVRTRGVGK..VPRKKVNVKVFI 241
 :|.:|.:|...||.:|...|...|.:|...|...|.:|...|.:|...| ||. .: .
 203 IGYVL..PLIIILICYSQICCKLFRTAQNPNLTEKSGVNKKALNTIILII 250
 :|.:|.:|...||.:|...|...|.:|...|...|.:|...|.:|...| ||. .: .
 242 IAVFFICFVPFHFARIPYTLSQTR..DVFDTAENTLFYVKESTLWLTS 289
 :|.:|.:|...||.:|...|...|.:|...|...|.:|...|.:|...| ||. .: .
 251 V.VFVLCTPTYHVIIQHMIKKLRFNSNLECSQRHSFQISLHFTVCLMNF 299
 :|.:|.:|...||.:|...|...|.:|...|...|.:|...|.:|...| ||. .: .
 290 NACLDPFIYFFLCKSFRNLSLISMILKCPNSATSLSQDNRKKEQDGDPNEE 339
 :|.:|.:|...||.:|...|...|.:|...|...|.:|...|.:|...| ||. .: .
 300 NCCMDPFIYFFACKGYKRKVMRMLKRQVS.VSISSAVKSAPEENSREMTE 348
 :|.:|.:|...||.:|...|...|.:|...|...|.:|...|.:|...| ||. .: .
 340 TPM 342
 :|.
 349 TQM 351

FIG.2

1	GGCACGAGCCCACCCCTGCGTCGGGCCTCAGTCAGCCCCGGGGAGGCCATGAACGCCAC	60
	M N A T	4
1		
61	GGGGACCCCGGTGGCCCCGAGTCCTGCCAACAGCTGGCGGCCGGCGGGCACAGCCGGCT	120
5	G T P V A P E S C Q Q L A A G G H S R L	24
121	CATTGTTCTGCACTACAACCACTCGGGCCGCTGGCCGGCGCGGGGGCCGGAGGATGG	180
25	I V L H Y N H S G R L A G R G G P E D G	44
181	CGGCCTGGGGCCCTGCGGGGCTGCGGTGGCCAGCTGCCTGGTGGCTGGAGAA	240
45	G L G A L R G L S V A A S C L V V L E N	64
241	CTTGCTGGTGCCTGGCGGCCATCACCAGCACATGCGGTGCGAACGCTGGGTCTACTATTG	300
65	L L V L A A I T S H M R S Q R W V Y Y C	84
301	CCTGGTGAACATTACGATGAGTGACCTGCTCACGGGCAGCGCCTACCTGGCCAACGTGCT	360
85	L V N I T M S D L L T G A A Y L A N V L	104
361	GCTGTCGGGGCCCGCACCTTCCGCTGGCGCCGCCAGTGGTCTACCGAAGGGCCT	420
105	L S G A R T F R L A P A Q W F L R K G L	124
421	GCTCTTCACCGCCCTGGCGCTCCACCTCAGCCTGCTTCACTGCAGGGTTGCGCTT	480
125	L F T A L A A S T F S L L F T A G L R F	144
481	TGCCACCATGGTGGCGCCGGTGGCGAGAGCGGGGCCACCAAGACCGAGCCGCTACGG	540
145	A T M V R P V A E S G A T K T S R V Y G	164
541	CTTCATGGCCTCTGGCTGGCTGGCCGCTGCTGGGGATGCTGCCTTGCTGGCTG	600
165	F I G L C W L L A A L L G M L P L L G W	184
601	GAACTGCGCTGCGCCTTGACCGCTGCTCCAGCCTTCTGCCCTACTCCAAGCGCTA	660
185	N C L C A F D R C S S L L P L Y S K R Y	204
661	CATCCTCTCTGCGCTGGTGATCTCGCCGGCTCTGGCCACCATCATGGCTATGG	720
205	I L F C L V I F A G V L A T I M G L Y G	224
721	GGCCATCTCCGCTGGTGAGGCCAGCGGGAGAAGGCCACGCCAGCGCCGGCCG	780
225	A I F R L V Q A S G Q K A P R P A A R R	244

FIG.3A

Appl. No. To be assigned; Group Art Unit: To be assigned
 Dkt. No. 1488.1220003/EXS/EJH
 Inventors: Li *et al.*; Tel: 202/371-2600
 Title: Human G-Protein Coupled Receptors

781	CAAGGCCCGCCGCTGCTGAAGACGGTGTATGATCCTGCTGGCCTTCTGGTGTGCTG	840
245	K A R R L L K T V L M I L L A F L V C W	264
841	GGGACCACTCTCGGGCTGCTGGCCGACGTCTTGGCTCCAACCTCTGGGCCAGGA	900
265	G P L F G L L A D V F G S N L W A Q E	284
901	GTACCTCGGGGATGGACTGGATCCTGGCCCTGGCGTCTCAACTCGGGTCAACCC	960
285	Y L R G M D W I L A L A V L N S A V N P	304
961	CATCATCTACTCTTCCGACAGCAGGGAGGTGTGCAGAGCCGTGCTCAGCTCCTCTGCTG	1020
305	I I Y S F R S R E V C R A V L S F L C C	324
1021	CGGGTGTCTCGGCTGGCATGCGAGGGCCGGGACTGCCTGGCCCGGGCGTCGAGGC	1080
325	G C L R L G M R G P G D C L A R A V E A	344
1081	TCACTCCGGAGCTTCCACCAACCGACAGCTCTGAGGCCAAGGGACAGCTTCGCGCTC	1140
345	H S G A S T T D S S L R P R D S F R G S	364
1141	CCGCTCGCTCAGCTTCCGGATGCGGGAGCCCTGTCCAGCATCTCCAGCGTGGAGCAT	1200
365	R S L S F R M R E P L S S I S S V R S I	384
1201	CTGAAGTTGAGTCTTGCCTGTGGATGGTCAACCACCGGGTGCCTGCCAGGCAGGCC	1260
385	*	385
1261	CCTGGGGTACAGGAAGCTGTGCAACGCAACCTGCCCTGATGGGGAGCAGGAACGGG	1320
1321	ACAGGCCCATGGACTTGCCTGGCTGCCAGCTCGGGCTCTGACGCCATATGGACTTG	1380
1381	CCATTGCCTATGGCTCACCTGGACAAGGAGGCAACCACCCACCTCCCCTAGGAGCAG	1440
1441	AGAGCACCCCTGGTGTGGGGCGAGTGGGCTCCACAACCCGCTCTGTTGATTCTGG	1500
1501	GGAAGTCCCGGCCCTCTGGGCTCAGTAGGGCTCCAGGCTGCAAGGGTGGACTGT	1560
1561	GGGATGCATGCCCTGGCAACATTGAAGTTGATCATGGTAAAAAAAAAAAAAAA	1620
1621	AAAAAAAAAAAAAAA	1637

FIG.3B

ppl. No. To be assigned; Group Art Unit: To be assigned
kt. No. 1488.1220003/EKS/EJH;
vendors: Li *et al.*; Tel: 202/371-2600
title: Human G-Protein Coupled Receptors

1	MNATGTPVAPESCQQLAAAGHSRLLVLYHYNHSGRLAGRGGPEDGGLGALR	50
	.. .:
1	MGPTSVPLVKAHRSSVSDYVNYDIIVRHYNYTGKLNISADKEN.SIKLTS	49
51	GLSVAASCLVVLENLLVLAITSHMRSQRWVYCLVNITMSDLTGAAYL	100
	.. .:	
50	VVFILICCFIILENIFVLLTIWTKKFHPRPMYFIGNIALSDLLAGVAYT	99
101	ANVLLSGARTFRLAPAQWFRLKGLLFTALAASTFSLLFTAGLRFATMVRP	150
	: :: . . : . . . :	
100	ANLLLSGATTYKLTPAQWFLREGSMFVALSASVFSLLAIAIERYITMLKM	149
151	VAESGATKTSRVYGFIGLCWLLAALLGMLPLLGWNCLCAFDRCSSLLPLY	200
 :: : :: : : : : 	
150	KLHNGS.NNFRFLLISACWVISLILGGLPIMGWNCISALSSCSTVLPLY	198
201	SKRYILFCLVIFAGVLATIMGLYGAIFRLVQASGQKAPRPAARRKARR..	248
	: 	
199	HKHYILFCFTVFTLLLSIVILYCRYISLVRTRSRLTFRKNISKASRSS	248
249LLKTVLMILLAFLVCWGPLFGLLLADVFGSNLWAQEYLRGMDWILA	294
	: 	
249	ENVALLKTVIIVLSVFIACWAPLFLILLLDV.GCKVKTCDILFRAEYFLV	297
295	LAVLNSAVNPPIIYSFRSREVCRAVLSFLCCGCLRLGMRGPDCLARAVEA	344
	: 	
298	LAVLNSGTNPPIYTLTNKEMRRAFIRIMSCCKCPSG..DSAGKFKRPIIA	345
345	...HSGASTTDSSLRPRDSFRGSRSLSFRMREPLSSS	379
	
346	GMEFSRSKSDNSHPKD..GDNPETIMSSGNVNSSS	381

FIG. 4